## In the Claims:

- 1-38. (Previously canceled).
- 39. (Presently amended) An isolated polypeptide having at least 80% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263, wherein said polypeptide induces chondrocyte redifferentiation is capable of inhibiting neoplastic cell growth.
- 40. (Presently amended) The isolated polypeptide of Claim 39 having at least 85% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263, wherein said polypeptide <u>induces chondrocyte redifferentiation</u> is capable of inhibiting neoplastic cell growth.

- 41. (Presently amended) The isolated polypeptide of Claim 39 having at least 90% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263, wherein said polypeptide induces chondrocyte redifferentiation is capable of inhibiting neoplastic cell growth.
- 42. (Presently amended) The isolated polypeptide of Claim 39 having at least 95% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263, wherein said polypeptide induces chondrocyte redifferentiation is capable of inhibiting neoplastic cell-growth.
- 43. (Presently amended) The isolated polypeptide of Claim 39 having at least 99% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);

- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide,
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263, wherein said polypeptide <u>induces chondrocyte redifferentiation</u> is capable of inhibiting neoplastic cell growth.
- 44. (Previously amended) An isolated polypeptide comprising:
- (a) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127);
- (b) the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127); or
- (d) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263.
- 45. (Previously added) The isolated polypeptide of Claim 44 comprising the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127).
- 46. (Previously added) The isolated polypeptide of Claim 44 comprising the amino acid sequence of the polypeptide shown in Figure 46 (SEQ ID NO:127), lacking its associated signal peptide.
- 47. (Previously added) The isolated polypeptide of Claim 44 comprising the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 46 (SEQ ID NO:127).

- 48. (Previously canceled).
- 49. (Previously added) The isolated polypeptide of Claim 44 comprising the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209263.
- 50. (Previously added) A chimeric polypeptide comprising a polypeptide according to Claim 39 fused to a heterologous polypeptide.
- 51. (Previously added) The chimeric polypeptide of Claim 50, wherein said heterologous polypeptide is an epitope tag or an Fc region of an immunoglobulin.